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BIOLOGICAL SCIENCES: Evolution

## **Termite queens close the sperm gates of eggs to switch from sexual to asexual reproduction**

**Short title: Queen control of egg fertilization in termites**

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**Males and females are in conflict over genetic transmission in the evolution of parthenogenesis because it enhances female reproductive output but deprives the males' genetic contribution. For males, any trait that coerces females into sexual reproduction should increase their fitness. However, in the termite *Reticulitermes speratus*, queens produce their replacements (neotenic queens) parthenogenetically while using normal sexual reproduction to produce other colony members. Here, we demonstrate that termite queens produce parthenogenetic offspring in the presence of kings by closing the micropyles (sperm gates; i.e., openings for sperm entry) of their eggs. Our field survey demonstrated that termite eggs show large variation in numbers of micropyles, with some having none. Microsatellite analysis demonstrated that embryos of micropyle-less eggs develop parthenogenetically, whereas those of eggs with micropyles are fertilized and develop sexually. Surveys of eggs among queens of different age groups showed that queens begin to lay micropyle-less eggs when they are older and thus need to produce their replacements parthenogenetically. In addition, we found clear seasonality in new neotenic queen differentiation and in micropyle-less egg production. This is the first identification of the mechanism through which females control egg fertilization over time in diploid animals, implying a novel route of the evolution of parthenogenesis in favor of female interests without interference from males.**

Keywords: Thelytoky, micropyle, asexual queen succession, Isoptera

## Significance

**To clarify the evolution of parthenogenesis, given the potential sexual conflict over genetic transmission, identifying the mechanism regulating egg fertilization in females is essential. In the termite *Reticulitermes speratus*, queens produce their replacements (neotenic queens) parthenogenetically, but use sexual reproduction to produce other colony members. We discovered that queens of the termite close micropyles (openings for sperm entry) of their eggs to produce parthenogenetic offspring in the presence of kings. Furthermore, we found that queens control the proportion of micropyle-less eggs by regulating of the number of micropyles over time. This study describes a novel route of the evolution of parthenogenesis in favor of females' interests without interference from males.**



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**T**he near ubiquity of sexual reproduction is one of the most enduring puzzles in evolutionary biology because, all else being equal, asexual populations have a twofold fitness advantage over their sexual counterparts and should rapidly outnumber sexual populations (1, 2). A proposed major advantage of sexual reproduction is that it promotes genetic variability across generations, facilitating adaptation to local ecological conditions (3–5). However, mathematical models have revealed that evolution need not favor sexual reproduction, even when it does increase variability that is beneficial, and thus the prevalence of sexual reproduction remains an enigma (6, 7). Some biologists have approached this question by considering how loss of sex can be achieved in a population of sexual organisms (8–10). The evolution of parthenogenesis may be no exception in that it creates sexual conflict, given that parthenogenetic reproduction enhances female reproductive output but deprives the males’ genetic contribution to future generations. Therefore, any trait in males that coerces parthenogenetic females into sexual reproduction should increase their fitness. A recent theoretical study demonstrated that the evolution of male coercion substantially favors the maintenance of sexual reproduction even though a female barrier against the coercion can evolve (10).

If male counteradaptations interfere with parthenogenesis, females would unlikely be able to switch from sexual to asexual reproduction unless isolated from males. Identifying the mechanism regulating egg fertilization is critical to better understand the evolutionary routes of parthenogenesis, given the potential sexual conflict involved. Unlike facultative parthenogenesis, whereby females use thelytoky only in the absence of males, queens of some ants (11–13) and termites (14–16) selectively use both sexual reproduction and thelytoky when mating with males.

In these social insects, queens produce new queens asexually by thelytokous parthenogenesis, but produce workers through sexual reproduction. This conditional switching of reproductive tactics allow queens to increase the transmission of their genes to the next generation and maintain genetic diversity in the worker force, but it inevitably provokes queen–male sexual conflict by reducing the males’ genetic contribution to future generations (17). This system provides an ideal opportunity to study how and why females switch from sexual to asexual reproduction even in the presence of males.

The breeding system of the subterranean termite *Reticulitermes speratus* is characterized by asexual queen succession (AQS) (Fig. 1 and *SI Text*). Termite colonies are typically founded by a monogamous pair of primary reproductives (adult winged forms), a king and queen. In *Reticulitermes* termites, primary queens live more than 11 years (18). As the primary queen senesces and thus her egg production becomes insufficient to maintain the colony, secondary queens (i.e., neotenic queens) differentiate within the colony and supplement egg production, eventually replacing the primary queen (Fig. 1 and Fig. S1) (18, 19). In AQS species, queens produce their neotenic replacement queens asexually, but use normal sexual reproduction to produce other colony members (14–16). Because sex determination in *Reticulitermes* termites is male heterogamy (20), queens cannot produce secondary kings by parthenogenesis (17). This AQS system enables founding queens to increase their reproductive output while retaining the same transmission rate of their genes to future generations. Therefore, the founder queen can be considered genetically immortal until the colony dies, as neotenic queens are also replaced by subsequent cohorts of parthenogenetically produced neotenic queens.

How can queens of termites (i.e., diploid social insects) control egg fertilization? In haplodiploid social Hymenoptera, unfertilized eggs become males and fertilized eggs produce

females (21), giving queens a potentially powerful mechanism for controlling fertilization (22, 23). However, in diploid insects, sperm release is generally activated through a neural loop whenever an egg passes the genital chamber, with no control over fertilization (24). In this study, we focused on micropyles of eggs to identify a possible mechanism for controlling fertilization in diploid insects. A micropyle is a channel that extends from the external gateway through the chorion and ends in the vitelline membrane, serving as the route of sperm entry into a mature oocyte (25–27). Termite queens are always attended by kings, and thus the simplest and most effective mechanism to produce parthenogenetic eggs would be to close the micropyles of eggs to prevent sperm entry. Preliminary observations of micropyles of the eggs of *R. speratus* showed that the number of micropyles varies largely among eggs, even within a colony. If a certain proportion of eggs are micropyle-less (i.e., eggs without sperm gates), fertilizing these eggs would be mechanically impossible, regardless of a king's presence. Unfertilized eggs of *R. speratus* restore diploidy through automixis with terminal fusion (28), and the embryos develop into parthenogenetic daughters (29). Thus, we hypothesized that queens control the production of parthenogenetic offspring independently of males by regulating the number of micropyles on their eggs.

We tested this hypothesis in a series of micropyle analyses and DNA genotyping of embryos aimed at confirming the production of micropyle-less eggs in field colonies and identifying the genetic profiles of eggs with and without micropyles. We also tested the likelihood of sperm depletion as a potential origin of parthenogenesis. We found that *R. speratus* queens alter the number of micropyles on their eggs over time and thereby produce micropyle-less eggs to asexually produce their replacement.

## Results and Discussion

To test the micropyle-less egg hypothesis, we first investigated the number of micropyles of eggs collected from 60 *R. speratus* field colonies. Micropyles were located on the posterior end of eggs, forming an arc (Fig. 2A). They are funnel-shaped with an opening of  $3.23\ \mu\text{m}$  ( $\pm 0.15$  SEM,  $n = 10$ ) in diameter (Fig. 2B), and the average number of micropyles for all samples ( $n = 6,000$ ) was  $9.48$  ( $\pm 0.04$  SEM, range = 0–33) (Fig. 2C and D). Significant differences were observed in the number of micropyles among colonies ( $F_{59, 5940} = 27.35$ ,  $P < 0.0001$ , one-way ANOVA), and seven of 60 colonies (11.7%) had micropyle-less eggs (Table S1). Unmated *R. speratus* queens lay eggs and the unfertilized eggs develop parthenogenetically, with a comparably high hatching rate as fertilized eggs (29). All micropyle-less eggs showed normal embryonic development as observed in previous studies on parthenogenesis in *R. speratus*. To test whether the embryos of micropyle-less eggs were developing parthenogenetically, we compared microsatellite genotypes of the embryos of eggs with and without micropyles in the four colonies (GB130502C, ZE130827B, KW140531A and AO140511A), which had micropyle-less eggs. In all of the colonies, the embryos of micropyle-less eggs had only maternal alleles, indicating parthenogenetic development, while those of eggs that had micropyles (even only a single micropyle) had both paternal and maternal alleles, indicating sexual development (Table 1 and Table S2). These results clearly indicate that the production of micropyle-less eggs functions as a mechanism for parthenogenetic reproduction by the queen in the presence of kings.

To investigate the occurrence of sperm depletion, we examined the sperm storage of secondary queens by dissecting out the spermathecae (sperm storage organs) and applying microsatellite genotyping. Collections of royal chambers from 54 field colonies revealed that

initially monogamous colonies became extremely polygynous as the colonies developed (Table S3). The number of queens per nest was  $57.63 (\pm 12.84 \text{ SEM}, n = 54)$ , and 51 of 54 colonies contained only a single primary king (Table S3). The largest colony (TA090620A) had 676 secondary queens and a single primary king. We detected sperm alleles (i.e., kings' alleles) from all sampled queens even in the colony containing 676 secondary queens (Tables S4–S6), suggesting that kings have the capacity to provide sufficient sperm, and thus parthenogenetic reproduction in colonies containing kings cannot be explained by sperm depletion.

For successful AQS, parthenogenetic daughters must exist in a colony whenever the colony requires supplementary or replacement queens. Conversely, overproduction of parthenogenetic offspring would impose a large cost on colonies because these individuals are inferior to sexually produced individuals as workers or alates, likely due to their complete loss of heterozygosity (14, 28). Therefore, selection should act against overproduction of parthenogenetic eggs and favor the production of necessary minimum numbers of micropyle-less eggs. If queens are able to control the production of micropyle-less eggs, we predicted that older queens that require asexual replacements would produce more micropyle-less eggs than young queens that will continue to reign for several years. We conducted seasonal sampling of queens from field colonies and identified the season of new secondary queen differentiation. In mid-May, just prior to swarming, parthenogenetic nymphs with small wing buds (pre-neotenic brachypterous stage; 30) develop into neotenic queens and sexual nymphs with long wing buds (alate-destined nymphs) molt into alates simultaneously (Fig. S2). Consequently, two types of secondary queens occur in the colonies under queen replacement in this season, old physogastric queens and first-year small queens, forming a clear bimodal size distribution (Fig. 3A). This seasonality of new secondary queen differentiation seems adaptive to maximize their contribution to egg production because

colonies begin to produce eggs in May and reach maximum output in July, followed by cessation in October (31). Among the 11 colonies, from which we could collect reproductives during the queen replacing season in 2013 and 2014, two colonies had new secondary reproductives but nine had only old physogastric queens, implying that secondary queen replacement occurs once in 5–6 years. We collected both types of queens from the two colonies (YO120508A and AO140511A) and compared the number of micropyles between eggs laid by old and young secondary queens. A significant difference was observed between the number of micropyles between the eggs laid by old secondary queens and those laid by young secondary queens (colony:  $F_{1, 396} = 105.93$ ,  $P < 0.0001$ ; age group:  $F_{1, 396} = 97.39$ ,  $P < 0.0001$ , two-way ANOVA), whereby the eggs laid by old secondary queens had significantly fewer micropyles than those laid by young secondary queens ( $P < 0.0001$ , Tukey's HSD). We also investigated the number of micropyles laid by young primary queens (foundresses) obtained from incipient colonies in the laboratory. A significant difference was observed in the number of micropyles between young primary queens and young secondary queens (colony:  $F_{3, 495} = 6.75$ ,  $P < 0.001$ ; queen type:  $F_{1, 495} = 10.24$ ,  $P < 0.01$ , nested-ANOVA), where the eggs laid by young primary queens had significantly larger number of micropyles than those laid by young secondary queens ( $P < 0.01$ , Tukey's HSD) (Fig. S3 and *SI Text*). Neither young primary queens nor young secondary queens produced micropyle-less eggs, suggesting that queens begin to lay parthenogenetic eggs to produce their asexual replacements when they are older. Why young primary queens do not produce micropyle-less eggs? Considering the high mortality of young queens in early founding stage, producing asexual offspring with the first brood might be adaptive for the queens. Importantly, workers of *Reticulitermes* termites retain totipotency to differentiate into ergatoid reproductives (worker-derived reproductives) (Fig. S1). Therefore,

workers develop into ergatoid queens and continue the colony if there is no parthenogenetic offspring in the colony, although it imposes the costs of inbreeding and reduces the queen's genetic contribution to the next generations from 1/2 to 1/4. Because the number of workers during early founding stage is crucial for colony survivorship, the cost of producing asexual offspring in the first brood would outweigh the cost of ergatoid queen replacements at the accidental death of primary queens. This can be the reason why we find no micropyle-less eggs among the eggs laid by young primary queens. In addition, micropyle-less egg production was observed almost exclusively in spring (Fig. 3B), suggesting that queens are also able to regulate the number of micropyles seasonally.

Our results confirm that queens of AQS termites control egg fertilization, and thereby regulate the proportion of sexual and asexual offspring. The proportion of parthenogenetic eggs  $P$  is determined by the cumulative distribution function:

$$P(\mu, \sigma^2) = \frac{1}{2} \left( 1 + \operatorname{erf} \frac{0 - \mu}{\sqrt{2\sigma^2}} \right),$$

where  $\mu$  ( $\mu \geq 0$ ) and  $\sigma^2$  are the median and variance, respectively, of the distribution of the number of micropyles (Fig. 3C). This indicates that parthenogenetic egg production increases as the median of the distribution decreases, and larger variance yields more micropyle-less eggs. Each micropyle is formed by a single cytoplasmic projection from a micropylar channel-forming cell (MCFC), and thus the variation in the number of the MCFCs is proximately responsible for the differing number of micropyles in insects (26, 27). Consequently, the expression of the gene that regulates the differentiation of MCFCs determines the switch between sexual and asexual reproduction of termites. It is known that the gene *hemipterous* (*hep*) is required in the follicle cells for morphogenesis of micropyles during oogenesis in *Drosophila* (32). Comparison of *hep*

expression between sexual and asexual strains in *Drosophila* species, such as *Drosophila mercatorum*, might illuminate the evolutionary linkage of parthenogenesis and micropyle formation in insects.

Considering the evolutionary process of parthenogenesis and AQS in termites, this switching mechanism based on the number of micropyles seems most parsimonious because reduction in the number of MCFCs as queens' age automatically results in the production of parthenogenetic offspring. AQS appears to have evolved in at least three *Reticulitermes* species independently; we would expect the other two AQS species, *R. virginicus* (15) and *R. lucifugus* (16), to have the same switching mechanism between sexual and asexual reproduction. The production of secondary queens through parthenogenesis effectively extends the reproductive life of the primary queen, greatly expanding her reproductive capacity. Coercion by kings cannot impede queens' parthenogenesis through micropyle-less egg production even though parthenogenesis reduces reproductive output by kings. This lack of counteradaptation by kings may have facilitated the evolution of AQS systems in termites.

In conclusion, this study demonstrated that queens of the termite *R. speratus* produce asexual offspring even in the presence of kings by laying micropyle-less eggs. The production of micropyle-less eggs gives females a powerful mechanism to control egg fertilization under sexual conflict over parthenogenesis. This study provides a mechanism for the evolution of parthenogenesis in favor of female interests independently of male interests.

## Methods



**Egg Collection from Field Colonies.** Sixty mature colonies of the subterranean termite *R. speratus* were collected from pine forests in Kyoto, Shiga, and Wakayama Prefecture, Kinki district, and Aomori Prefecture, Tohoku district, Japan, from May 2012 to May 2014. All sampling was conducted from May to October (Table S1) because a previous study revealed a seasonal pattern of egg production in *R. speratus*—colonies begin to produce eggs in May and reach the maximum level in July, followed by cessation in October (31). We very carefully dismantled nest wood and extracted eggs using an aspirator. The reproductives (kings and queens) were also collected if they were present. The eggs and reproductives collected from each colony were placed in a moist unwoven cloth in a 90-mm Petri dish together with nursing workers and transported to the laboratory.

**Micropyle Analysis.** We investigated the micropyle morphology and distribution on eggs of *R. speratus* using a scanning electron microscope (VE-8800; Keyence) and a laser 3D measurement microscope (VK-X200; Keyence). Size measurements of 10 eggs randomly selected from the eggs extracted from field colony OO140529B (Kyoto) were obtained using the scanning electron microscope.

We developed a micropyle staining method to observe micropyles without damaging DNA. Live eggs collected from each colony were placed in a sterile 1.5-mL tube containing 500  $\mu$ L of 1% eosin Y solution (Wako Pure Chemical Industries) and kept for 3 h at room temperature before observation. Eggs preserved in eosin Y solution can be used for micropyle observation and for microsatellite analysis for 2 weeks if kept at 4°C. Each focal egg was placed on a moist unwoven cloth in a 35-mm Petri dish in a posterior-up position using a stereomicroscope (SZX7; Olympus), and the number of micropyles was counted under a digital microscope (VHX-900;

Keyence). One hundred eggs were randomly chosen from each colony (6,000 eggs in total) to determine the number of micropyles, and these data were analyzed using a one-way ANOVA. To determine the seasonal pattern of micropyle-less egg production, we also compared the proportion of colonies with and without micropyle-less eggs among the colonies collected in May (spring), from June to August (summer), and from September to October (autumn) using Fisher's exact probability tests with sequential Bonferroni correction.

**Microsatellite DNA Genotyping.** We used four field colonies collected in Wakayama (colony GB130502C), Shiga (ZE130827B), Kyoto (KW140531A) and Aomori (AO140511A) for microsatellite analysis of embryos. After micropyle analysis, we sorted the eggs based on the number of micropyles; three eggs were randomly chosen from each egg group with 0, 1, 2, 3, 4, or 5 micropyles for each colony (Table 1 and Table S2). Eggs with embryos of developmental stage II or later (29) were used for microsatellite analysis to exclude any non-developing eggs. We also genotyped the single primary king and 10 randomly chosen secondary queens from each colony to assign parentage of the eggs (see Table S3 for the royal composition of each colony). All individuals used in this analysis were placed in vials containing 99.5% ethanol and stored until DNA extraction. Whole eggs or heads of individual termites were ground in Chelex-100 resin solution (Bio-Rad), and DNA was extracted and purified in accordance with standard Chelex-based protocols (14). Individuals were genotyped at five highly polymorphic microsatellite loci: *Rf 6-1*, *Rf 21-1*, *Rf 24-2* (33), *Rs 10*, and *Rs 15* (34). PCR conditions are detailed in a previous study (33, 34); fluorescently labeled PCR products were analyzed in a 3500 Genetic Analyzer (Applied Biosystems) with the internal GeneScan-600 LIZ size standard (Applied Biosystems). Allele sizes were determined using GeneMapper 5 (Applied Biosystems).

Embryos were determined as sexual (containing both paternal and maternal alleles) or asexual offspring (only containing maternal alleles) based on the genotypes of the five microsatellite loci.

**Micropyle Comparison between Old and New Queens.** Differentiation of secondary queens (neotenic queens) occurs simultaneously in mid-May just prior to swarming (Fig. S2).

Consequently, two types of secondary queens occur in the colonies under queen replacement in this season: the first-year small queens (2.5–4.1 mg fresh weight) and old physogastric queens (5.5–8.3 mg fresh weight), forming a clear bimodal size distribution (Fig. 3A). To compare the number of micropyles between the queen age groups, we used two field colonies (YO120508A and AO140511A), which had both types of queens, collected in Kyoto and Aomori, respectively. The new and old secondary queens were separated from each colony and placed on a moist unwoven cloth in a 90-mm Petri dish together with 100 workers. The Petri dishes were kept at 25°C for 1 week to obtain the eggs of the queens of each age group. One hundred eggs laid by the queens were randomly chosen from each age group, and we counted the number of micropyles using a digital microscope (VHX-900; Keyence) after staining. The numbers of micropyles of eggs were analyzed using a two-way ANOVA followed by Tukey's HSD test.

We also examined the number of micropyles of the eggs laid by young primary queens. The nests of three *R. speratus* colonies (UR140513F, OO140531A and OO140603A) were collected in Kyoto, Japan just before swarming in 2014. After alates emerged from the nest, they were separated by sex and maintained in Petri dishes containing moist filter paper until they shed their wings. Then, a male and a female were randomly chosen from each colony and placed in a 90-mm Petri dish that contained mixed sawdust bait blocks. Eight replications were made for each colony. The Petri dishes were kept at 25°C under constant darkness. After 35 days, the nests in

bait blocks were dissected to collect eggs from the incipient colonies. Because the number of eggs in each incipient colony was limited (9–19 eggs), one hundred eggs were randomly chosen from the eight incipient colonies, whose founding pairs were originated from the same colony. To compare the numbers of micropyles between the eggs laid by young primary queens and those laid by young secondary queens, we used a two-way nested ANOVA followed by Tukey's HSD test.

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## Figure legends

**Fig. 1.** Colony development and asexual queen succession in the termite *Reticulitermes speratus*.

As the primary queen senesces, secondary queens produced asexually by the primary queen differentiate within the colony and supplement egg production, eventually replacing the primary queen (details described in *SI Text*). PK, primary king; PQ, primary queen; SQ, secondary queen. Squares indicate males and circles represent females. Scale bars, 3 mm.

**Fig. 2.** Variation in the number of micropyles of eggs. (A) Scanning electron microscope image of the posterior end of an egg with micropyles (ventral view). Scale bar, 50  $\mu$ m. (B) Confocal scanning laser microscope image of micropyles (close-up view). Scale bar, 10  $\mu$ m. (C) Posterior views (dorsal up) of representative eggs with different number of micropyles (No. MP: 0, 2, 4, and 9). Micropyles were visualized by eosin Y staining. Scale bar, 100  $\mu$ m. (D) Comparison of the number of micropyles among four representative colonies ( $n = 100$  for each colony) and the total of the 60 field colonies ( $n = 6,000$ ). Micropyle-less eggs are indicated by red bars.

**Fig. 3.** Regulation of the production of micropyle-less eggs by the queens. (A) Comparison of the size of newly differentiated secondary queens (young SQ) and old physogastric secondary queens (old SQ) (upper panels) in colonies YO120508A (left) and AO140511A (right), and frequency distributions of the number of micropyles of the eggs laid by young (middle panels) and old queens (lower panels). Micropyle-less eggs are indicated by red bars. Blue lines show the fitted normal distributions. (B) Comparison of the proportion of the colonies with (closed bars) and without micropyle-less eggs (open bars) among the colonies collected in spring ( $n = 9$ ), summer ( $n = 35$ ), and autumn ( $n = 16$ ). Different letters on the bars indicate significant

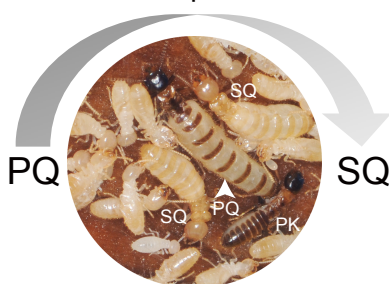
differences ( $P < 0.01$ , Fisher's exact probability test with sequential Bonferroni correction). (C)

Expectation of the proportion of micropyle-less eggs based on the median and the variance of the distribution of the number of micropyles.

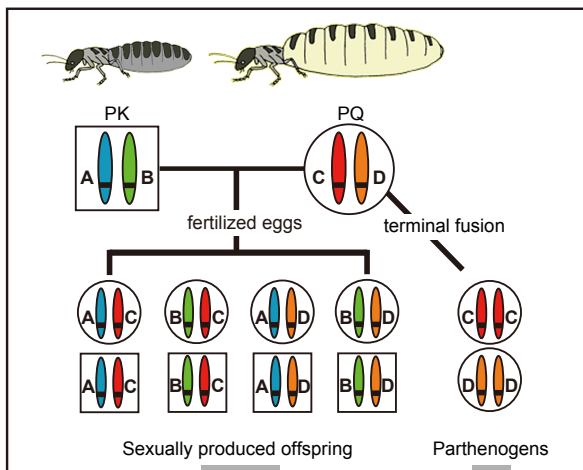
Incipient colony



Queen replacement

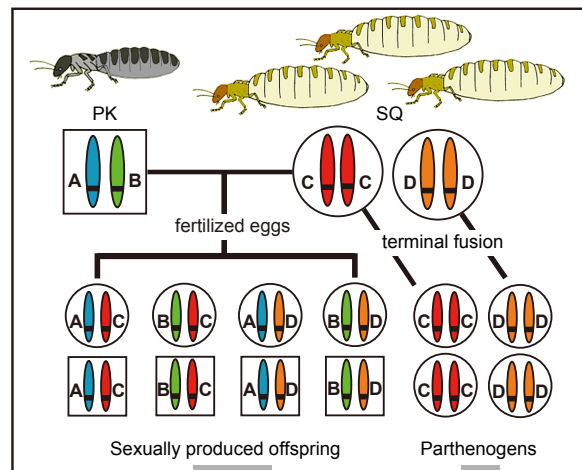


Mature colony



Differentiation into  
alates and workers

Differentiation into  
secondary queens



Differentiation into  
alates and workers

Differentiation into  
secondary queens

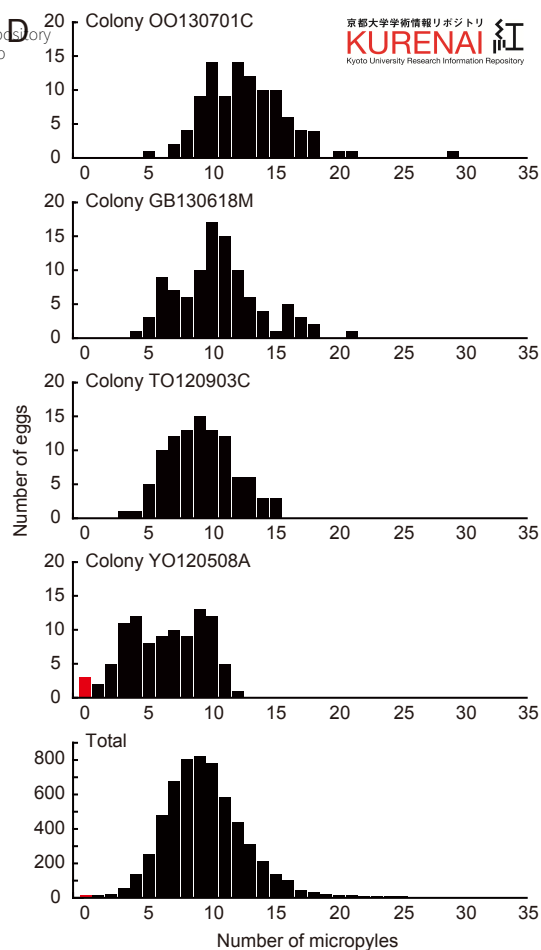
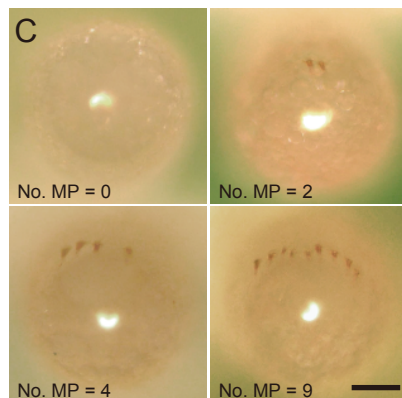
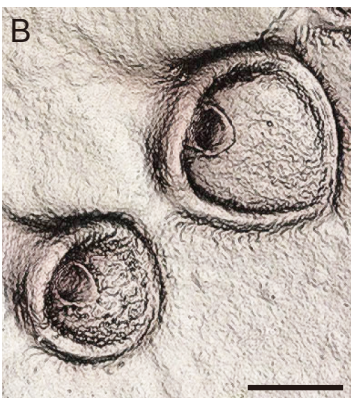
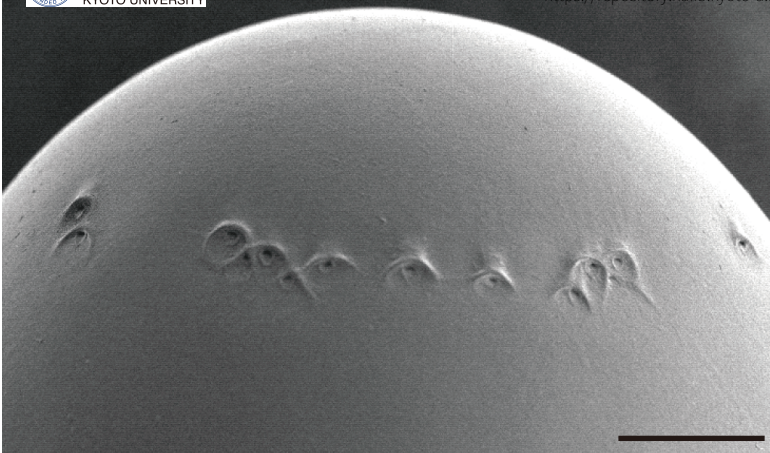


Fig. 2

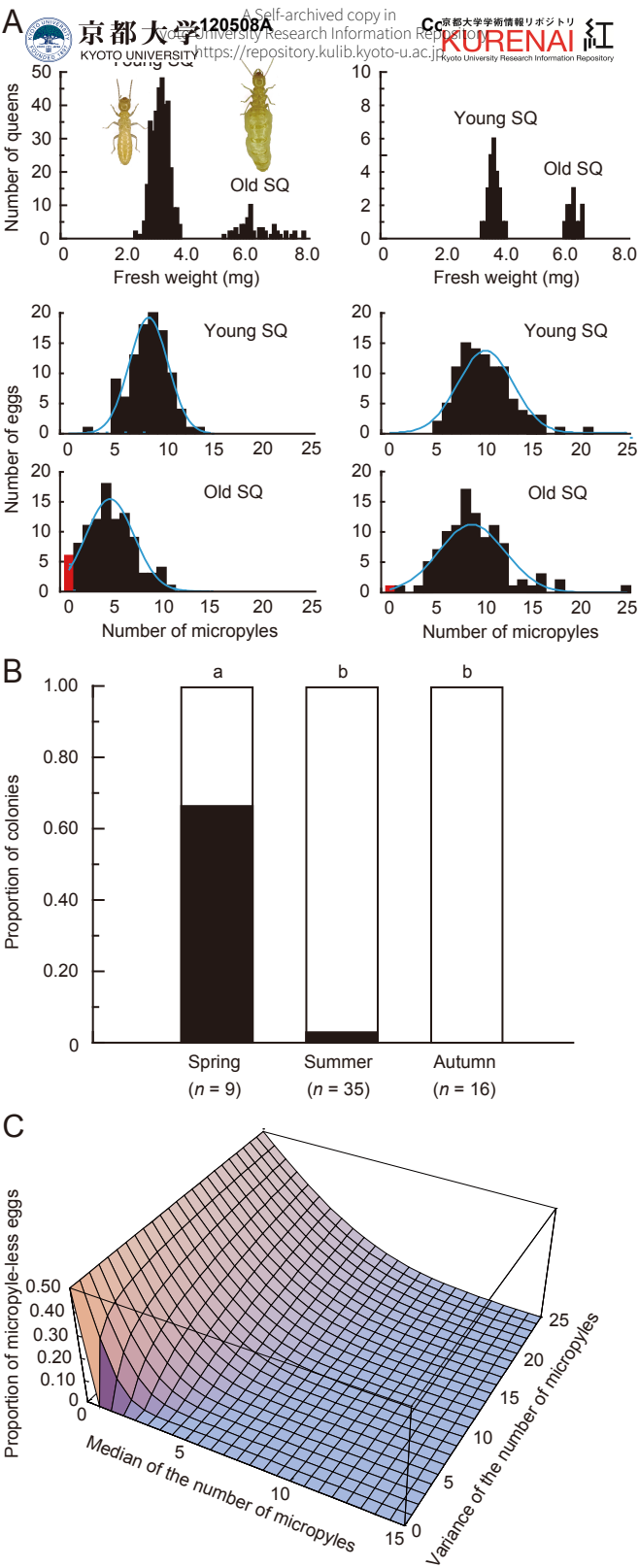


Fig. 3

Table 1. Genotypes of embryos (E) of the eggs with or without micropyles (MP), primary kings (PK), primary queens (PQ), and secondary (neotenic) queens (SQ) in colony GB130502C and ZE130827B at each of the five microsatellite loci.

Colony GB130502C							Colony ZE130827B						
Individual	Rf 24-2	Rf 21-1	Rs 10	Rs 15	Rf 6-1	P/S*	Individual	Rf 24-2	Rf 21-1	Rs 10	Rs 15	Rf 6-1	P/S*
Embryo (No. MP)							Embryo (No. MP)						
E-1 (0)	99 99	202 202	152 152	291 291	161 161	P	E-1 (0)	90 90	226 226	140 140	270 270	167 167	P
E-2 (0)	102 102	196 196	152 152	291 291	161 161	P	E-2 (0)	90 90	223 223	140 140	270 270	152 152	P
E-3 (0)	102 102	196 196	152 152	291 291	161 161	P	E-3 (0)	90 90	223 223	140 140	255 255	152 152	P
E-4 (1)	99 87	241 202	155 152	291 279	161 161	S	E-4 (1)	114 90	226 205	140 140	273 270	167 167	S
E-5 (1)	102 87	265 196	161 152	291 273	161 161	S	E-5 (1)	105 87	226 205	140 140	273 255	167 152	S
E-6 (1)	102 99	265 196	155 152	291 279	161 152	S	E-6 (1)	90 87	226 205	188 164	270 270	167 167	S
E-7 (2)	102 87	265 196	161 152	291 273	161 152	S	E-7 (2)	90 87	226 223	188 164	270 270	167 167	S
E-8 (2)	102 99	241 196	155 152	291 279	161 152	S	E-8 (2)	90 87	226 205	188 164	270 270	167 167	S
E-9 (2)	102 87	241 196	161 152	291 273	161 152	S	E-9 (2)	114 90	226 205	140 140	273 270	167 167	S
E-10 (3)	102 99	241 202	161 152	291 273	170 161	S	E-10 (3)	114 105	223 205	188 164	273 255	167 167	S
E-11 (3)	102 99	241 196	161 140	291 273	161 161	S	E-11 (3)	114 90	226 223	140 140	270 270	167 152	S
E-12 (3)	102 99	241 196	155 152	291 279	161 161	S	E-12 (3)	90 87	223 223	140 140	273 270	167 152	S
E-13 (4)	102 87	241 196	161 152	291 273	161 152	S	E-13 (4)	114 105	223 205	188 164	270 270	167 167	S
E-14 (4)	102 99	265 202	161 152	291 273	161 152	S	E-14 (4)	114 90	223 205	140 140	273 255	167 152	S
E-15 (4)	99 87	265 202	161 152	291 273	161 152	S	E-15 (4)	114 90	223 205	140 140	270 255	167 152	S
E-16 (5)	102 99	265 196	161 140	291 273	161 152	S	E-16 (5)	105 87	223 205	140 140	270 270	167 152	S
E-17 (5)	99 87	241 202	161 140	291 273	161 161	S	E-17 (5)	114 90	223 205	188 140	270 270	167 152	S
E-18 (5)	102 87	241 196	161 140	291 273	161 161	S	E-18 (5)	114 90	223 205	140 140	273 255	167 152	S
PK	102 87	265 241	161 155	279 273	161 152		PK	114 87	223 205	188 140	273 270	167 167	
PQ†	102 99	202 196	152 140	291 291	170 161		PQ†	105 90	226 223	164 140	270 255	167 152	
SQ							SQ						
SQ-1	102 102	196 196	152 152	291 291	161 161		SQ-1	105 105	223 223	164 164	270 270	167 167	
SQ-2	102 102	196 196	152 152	291 291	161 161		SQ-2	105 105	223 223	140 140	270 270	152 152	
SQ-3	102 102	196 196	140 140	291 291	161 161		SQ-3	90 90	223 223	140 140	270 270	152 152	
SQ-4	102 102	202 202	140 140	291 291	170 170		SQ-4	105 105	226 226	140 140	255 255	152 152	
SQ-5	99 99	202 202	140 140	291 291	161 161		SQ-5	105 105	223 223	140 140	270 270	152 152	
SQ-6	102 102	196 196	140 140	291 291	161 161		SQ-6	90 90	226 226	140 140	270 270	167 167	
SQ-7	99 99	196 196	140 140	291 291	170 170		SQ-7	105 105	226 226	140 140	270 270	152 152	
SQ-8	99 99	202 202	152 152	291 291	161 161		SQ-8	105 105	226 226	140 140	270 270	152 152	
SQ-9	99 99	196 196	140 140	291 291	161 161		SQ-9	90 90	223 223	140 140	270 270	152 152	
SQ-10	102 102	196 196	152 152	291 291	161 161		SQ-10	105 105	226 226	140 140	270 270	152 152	

Maternal alleles are indicated in red, and paternal are in blue.  
\*P/S: Parthenogenetically developed (P) or sexually developed (S).  
†The genotype of primary queens was determined from the genotype of offspring since primary queens had been replaced by parthenogenetically produced secondary queens.